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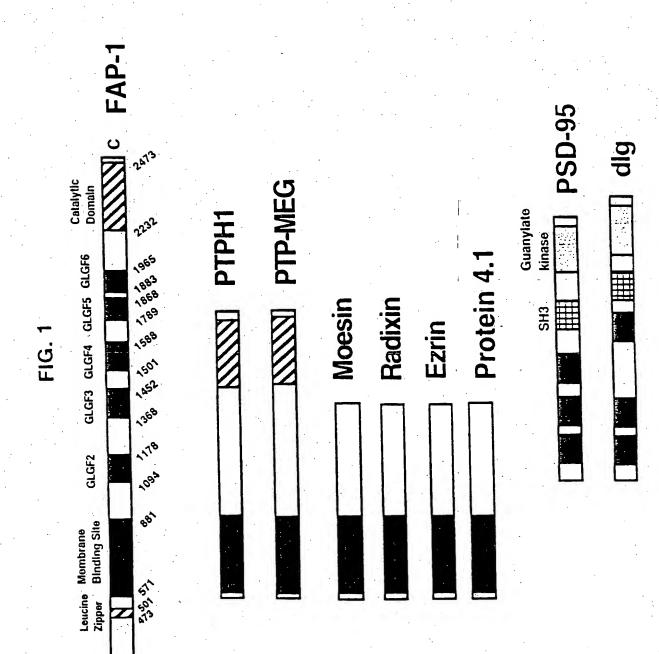
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2

GLGF1

Construction of pBTM116 (LexA)-(X)15 DUP16-FRP-1 Library DNAs of pBTM116 (LexA)-(X)15 Large scale transformation of yeast L40 CEHR-(H)15 PUP16-FRP-1 His+,  $\beta$ -gal+ Curing of pVP16-FAP-1 Isolation of

Analysis of DNA sequences

pBTM116 (LexA)-(X)15

8-1 9-3

0-5

14-1

•	•	
DSENSNFRNEIOSLV	SISNSRNENEGOSLE	STPDTGNENEGOCLE
lG. 28 Human	Rat	Mouse

FIG. 2C

FIG. 2D

VISOBERS	5   ·	ш	z	-	NMNEL	T Y   V		RGFISSLV	> Ø	
C	0-71	2-0	13-0	20-0	6-2	9-2	18-1	22-1	71-1	14-5
	C Y A   A  G   L   V	ENA GVS E	WWG ATO P	E H A Q Q Q	NSS FHS L	GLR LPP D	GSD SGV N	X K K N N N X	I G K D V W A	ASR NEE L I

25-9

57-5 72-1

16-13

### FIG. 3A

Fas C terminal 15 a.a. peptide ( $\mu$ M)



200 -

97.4 - 69 -

FAP-1

46 -

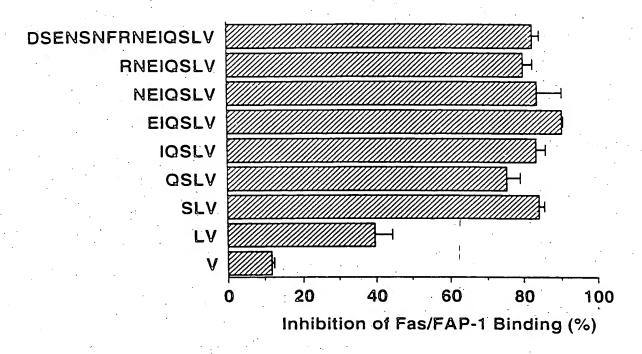
30 -

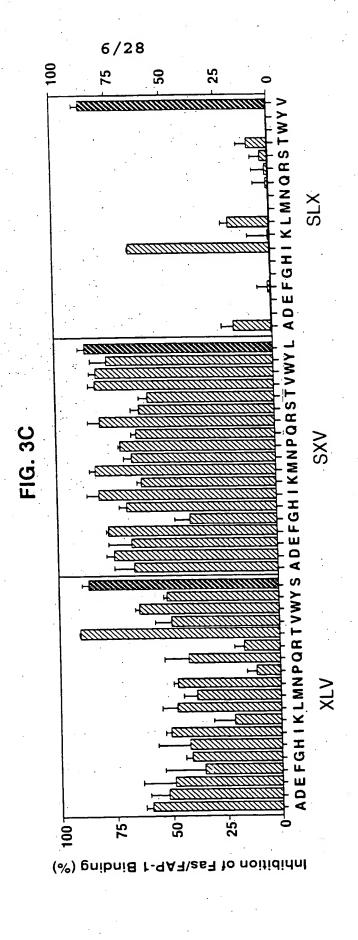
21.5 -

14.3 -

1 2 3 4 5 6 7 8 9 10

FIG. 3B





### FIG. 4A

VP	16	VP16	3
FAP-1	Ras	FAP-1	Ras
	000		
) O E	000		The St. C.
		C 40 40	
	000	THE E	6 G E
CO	066	200	# F 3
	FAP-1		FAP-1 Ras FAP-1

His +

His -

FIG. 4B



250 -148 -

← FAP-1

60 -

42 -

30 -

1 2 3

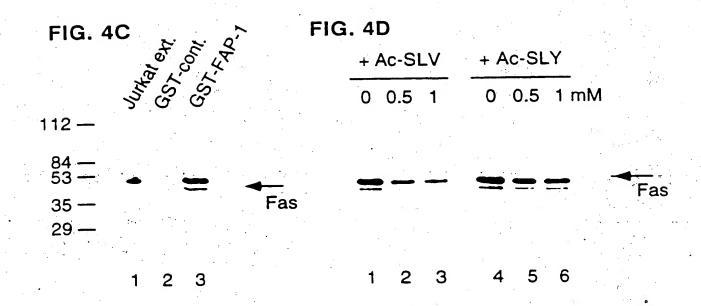
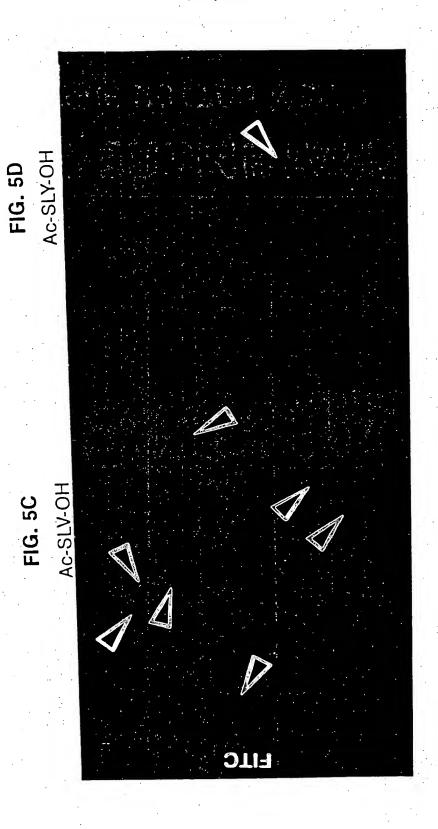




FIG. 5A

FIG. 5B



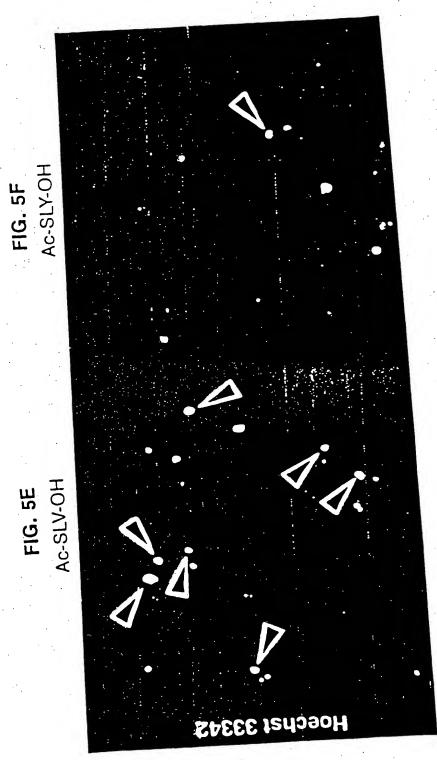
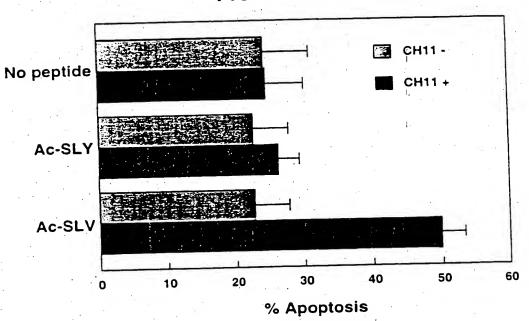


FIG. 5E





### FIG. 7A

Receptor

NGF

edterglrec vvttvmgssq nqtpppegek Ingsagdtwr adlveslcse ygyyddettg egvaqpcgan eaddavcrca qdliastvag nkqgansrpv gecckacnlg pakreevekl vdpclpctvc llaalrrigr acptglyths gldsmsapcv dgtysdeanh ddsslysslp afkrwnsckg stdepeappe atqdsatlda lgvslggake epckpctecv cpvrallasw kgntvceecp ppegsdstap tqtasgqalk avvvglvayi lipvycsila sqslhdqqph ehidsfthea gsglvfscqd ipgrwitrst dgprllllll vtfsdvvsat pvvtrgttdn mgagatgram qtvcepclds **lhsdsgisvd** hlagelgyqp trwadaecee rceacrycea stat**spv** 421 361 181 241 301

### FIG. 7B

### CD4 Receptor

yagsgnltla kltgsgelww vskrekavwv vsqlelqdsg figlgiffcv fhwknsngik vedgkeevgl iedsdtyice sfplaftve hltlpgalpg ctasqkksiq kniqggktls slklenkeak vlggvaglll llvlqlallp aatqgkkvvl gkkgdtvelt dsrrslwdgg nfpliiknlk vykkegegve stpvqpmali psydcrsprg klqmgkklpl qlqknltcev wgptspklml fqktc**spi** ekktcdcphr vsvkrvtqdp esnikvlptw ltlesppgss vlafqkassi rmsgikrlls kgpsklndra kkvefkidiv witfdlknke thllqgqslt evnlvvmrat 11sdsgqv11rcrhrrrgae **ewtctvlqnq** gaerasssks Inpeagmwdc ilgnqgsflt leaktgklhg lvfgltansd mnrgvpfrhl 41 361

FIG. 7C

Species	C-terminal sequences of NGFR (p75)	Binding activity of FAP-1
Human	SESTATSPV-COOH	+
Rat	SESTATSPV-COOH	+
Chicken	SESTATSPV-COOH	+

### FIG. 70

rttcsenela agesvqpwes nlvaayekak wekelagire gpsspgrits lesigvsssv sibidplsy streageday kklakaqceq selrselsgs nvvcgrkkss slilgqfraa csqmd]lt1 dgecggafav fundlkrans qtererdlle ssdrpvlgse llalaesed elgrvitgle gtrlqsvqat ypnlaeersr eringrichi kekkalelkl gttireedey aspalelael aavkitmlel rialleeens seirhqqsae ysagcieaye dyiqq1\mdr sikaqiyile dkpgkecada divelnkrlq elnkkidrlg dadacsdins clsktreess skirefevet naakallmkl elegylgrdl drlrrrvrel vsaleritks slsstssgsk lyshgsalse rahdcrktae elmamkeema .plakiaerv natalrlalq ftkedegr1k hcdlaiktve ritelhsvia pengetmyta hsaalaslkg armaireer esquanter gdenitomlk csniqeifqt memlvgkyee sstasscdte qerttlryee hieglttase ldlenavlmq veedkagrmr kklkarvqel elatssssnd mddddtsvs1 skaaalnrtk ndssaelsel aeftnairre lvhiehlkse ach ahslqd rygsepgdas senahtstt lvkprgdsgr enesitamic nrpinpstg shlmrehedv ghevnedsrs eecksnaer mnsgvamkyg cslsvaevdr 661 301 361 421 601 181 241 481 241

### FIG. 7E

pegdeegrame gkvmladrkg mlagqppfdg drlyfvmeyv ldseghikia vrehaffrri sdfeafsyvn shctdfiwgf sptfcdhcgs eklhvtvrda kpsdkdrrl wwaygvllye dalvianidq lgcgpegerd arffkoptfc riylkaevad pownesftfk Lavigkgst calhecfatv yrdlkldnvm chkfkihtyg egeyynvpi ergapy1tpp kgpdtddprs ayqpygkavd glmtkbpakr candhtekrg ktktirstin vhevkdhkf1 sgwykllng drvkltdfn allddopfl. Lfflhkrgii vtfscpgad pdpkneskq Egvs@lmkmp vectimekry vfyaaeisig pdyiapeii. skeavsick kgaenfdkff rkgalrdm sedrkapsnn kgcvinvpsl ckdwigddd mehavsypks tradfmgsls gpagnkvisp qvgkfkepqa **appfkpkvcg** dgvttrtfcg cfvvhkrche **tasqdvanr** f cetechmyh sdpyvk1k1 dweklenre1 adadelfqsi madvfpgnds gkageacave nggdlmybiq [fgmckehum lyglihqgm in 1 ipmdpng sveiwdwdrt eelyaikil rokfekak

### FIG. 7F

[dryvaignp seklfgrsih escnedvíga vntipalayk yflmslaiad ddnfvligef nrtalscedo edgvnekv**eg** v enkkplqlil itrimavick vfkegsclla dafnwtvdse lekklqnatn asimhlcais stlpqsele qhseeaskdn fsrylqcqyk gnilvimavs pvfglgddak digtraklas 1fvvmwcpff wiyldvlfst dfnsgeants tisvgismpi lqkeatlevs ndcsmvalgk tlfnktyrsa tavviiltia wplpsklcav ackvlgivff lnddtrlysn lssavnplvy vityfltiks tmgs 1 sneqk kaflkilavw smitilygyr lestrnslmg 1qekmwsall lspsclallh mllgflympv vsffipleim 11nvfvwigy cepgaytgrr seglongakk md:1ceents hinsrfngrt

### FIG. 7G

gdkteeqwex dscnqttlgm reakiyfrnp aímivtyflt kalpnøgdet cafikituw alltinfeam gnklhwaall dtllltene atkevktlrk eemkqiveed llvglfvmpf lgangynera *r*amldgsrkd fithitivic slaafftpl wagldtesip yflmelavad vdryiaikkp ker fgdfml E detpcsspak Elflimopf fqryitemyr npnniccv1t witvstvfqr raskvlgivf tl£nktfrda lekklqyatn asimhleais stfvhvissn gmtlvilavs wlfldvlfst hgirnginpa pikgietdvd vknkppqrlt vasgvnplvy ksvatiened isigiaipv maenskffkk wplplvlcpa mrrtstigk Lleifwigy ilmvilptig halqkkayl malsyrvse 181 241 301 361 421

### FIG. 7H

```
i maaasydqii kqvealkmen snirqeledn snhitklete asnmkevikq iqqsiedeam
 il assggidle rikelnidss nipgvklrsk msirsygere gsvssrsged apvpngsipr
121 rgfvngsres tgyleeleka rsliladidk eekekdwyya qiqmitkrid slpltenfsi
181 qtdmtrrqle yearqirvam eeqlqtcqdm ekraqrriar iqqiekdilr irqllqsqat
241 eaerssqukh eigshdaerg negggvçein matagngggs tirmdhetas vissssthaa
301 prritshigt kvemvyslis migthdkddm srtliamses gdscismrqs golpiliqli
361 hgndkdsvli gmsrgskear arasaalhni ihsopddkrg rreirvinii eqiraycetc
421 wewgeahepg mdqddcnpmpa pvehgicpay cylmkiside enrhamnelg glqaiaellq
461 vdcemygltm dhysitlrry agmaltnitf gdvankatle smkgemralv açıksesedi
541 qqviasvlrn lswradvnsk ktlrevgsvk almecalevk kestlksvls alwnisahc:
601 enkadicavd galaflygtl tyrsqtntla iiesgggilr avssliatne dhrqilrenn
661 clqtllqhlk shsltivsna cqtlwnlsar npkdqealwd mgavsmlknl ihskhicmiam
721 gsaaairnim anrpakykda nimspgssip sihvrkqkal eaeldaqhis etfdnichis
781 pkashrskqr hkqslygdyv fdtnrhddnr sdnfntgnmt vispylnttv lpsssssrqs
 841 ldasraekdr slerergigl gnyhpatenp gtaskrolqi sttaaqiakv meevsaihta
901 qedrssgatt elhovtdern alrrasaaht hantynitka eranrtosmo yakleykras
961 ndslnavses dgygkrgqmk psiesysedd eakfosygdy padlabkihs arbmddndge
1021 ldcpinyslk ysdeglnagr gspagnerwa rpkhiledei kqaeqrqarn qsttypvyte
1081 stddkhlkfq phfgqqecvs pyrsrgangs etnrvgenig inqnvsqslc qeddyeddkp
1141 thyserysee eqheeerpt nysikyneek rhvdqpidys lkyatdipss qkqsfsfsks
1201 saggsakteh messsentst pasnakrong lhpssagsrs gopokaatek vasingetig
1261 tycvedtpic ferceslasi seaedeigen qttqeadean tiqiaeikek igtreaedpv
1321 sevpaysqhp rikssrlqgs slssesarhk avefssgaks paksgaqtpk sppehyvqet
1381 plmfarctsv ssldsfesrs lasavgsepc sgmvagilap adlpdapggt mppsraktpp
1441 pppgtagtkr evpknkapta ekresgpkga avnaavgrvg vlpdadtilh fatestpdgf
1501 scassisals idepfickdy elrimppyge ndngmetase opkesnenge keaektidse
1561 kdilddøddd dieileecii samptkeerk akkpaqtask lpppvarkps glpvyklips
1621 qurlqpqkhv sftpqddmpr vycvegtpin fstatsledi tiesppnela agegvrggaq
1681 sgefekrdti ptegrstdea gggktssvti pelddnkaee gdilaecins ampkgkshkp
1741 frykkindav agasessap næglagkkk ketspykpip anteyrtryr knadskanla
1801 aervisdika skkonlkons kainaklpon edrvigsiai aspinytpie gtpycisma
1861 slasldfddd dydlsrekae irkakenkes eakytshtel tangqsankt qalakqpinr
 1921 gapkpilaka stipasski pargaatdek lanfalentp voishnssls sladidaenn
 1981 nkenepiket eppdaggeps kpqasgyapk sfhvedtpvc fsrasslssi sidseddllq
 2041 ecissampkk kkpsrlkgdn ekhspringg ilgeditldl kdigipdseh glspdsenfd
 2101 wkaigegans ivssihqaaa aacisrqass dsdsiislks gislgspfhl tpdqeexpft
 2161 ankaprilkp gekstletkk ieseskgikg gkkvykslit gkvrsnseis ggmkaplaan
 2221 mpsisrgrtm ihipgvrnss sstspyskkg pplktpasks psegqtatts prgakpsyks
 2231 elapvarque qiggeskape regerdetpe rpacquelerp iceporneis pgrngispon
 2341 klsqlprtss patastkssg sgkmaytspg romsqqrltk qtglskmass iprsesaskg
 2401 lnqmngnga nkkvelsrms stkssgsesd rserpvlvrq stfikeapsp tlrrkleesa
 2461 efeslapest pasptraged tpvlspslpd malathasvq aggwrklppm laptisyndg
 2521 rpakrhdiar shaesparly immagtwkre hakhaaslpr vatwrrtgaa asilsasses
2531 sekaksedek hynaisgtko skenovako twrkikenef aptnatagtv asgatngaes
 2541 ktliyqmapa vsktedvwvr iedopinnpr sgrsptgntp pvidsvseka npnikdskdn
 2701 qakqnvgngs vpmrtvglen rlnsfiqvda pdqkgteikp gqnnpvpvse thessivert
 2761 pissesskh espegivaar vipinynpep rkssadsisa rpsqipipvm nnikkrdekt
 2621 datessgtqs pkrhagsylv ter
```

FIG. 8

## (Low-affinity nerve growth factor receptor) p75NGFR

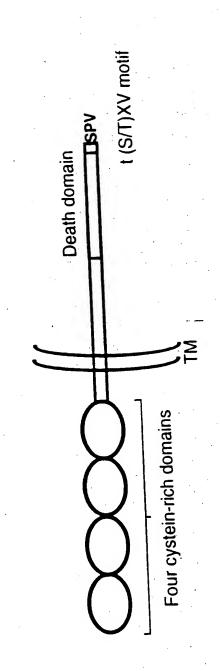


FIG. 9

C-terminal amino acid sequence	NEIQSLV	STATSPV
	Fas	p75NGFR

PDZ domain t (S/T)-X-V - COOH

interaction

FIG. 10

# In vitro interaction of 35S-labeled FAP-1 with various receptors FAP-1 binds to the cytoplasmic region of p75NGFR.

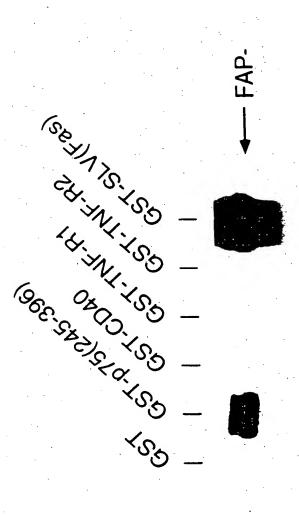
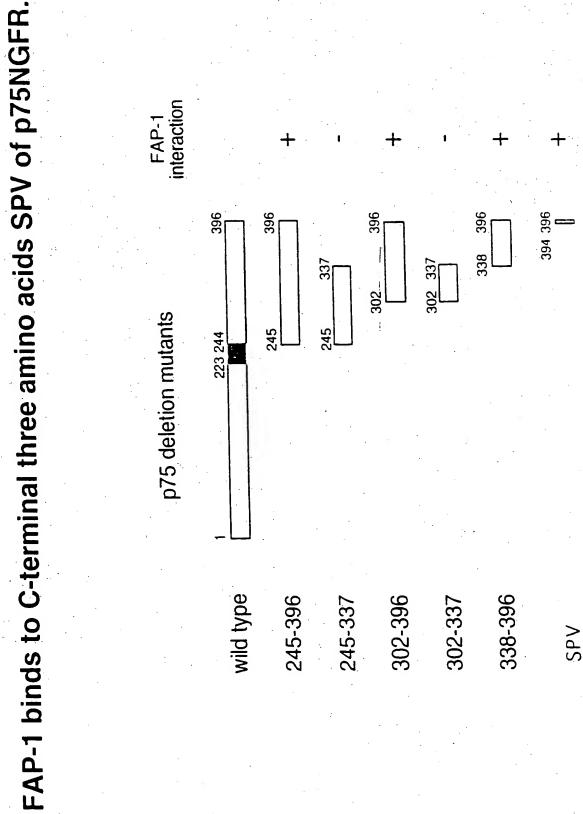


FIG. 11A



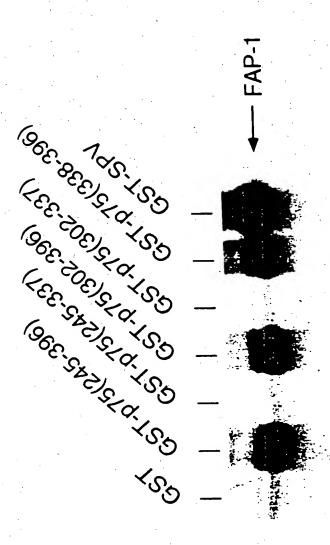


FIG. 11B

FIG. 12

# FAP-1 binds to p75NGFR C-terminal cytoplasmic region in yeast

	VP16-FAP-1	VP16-cRaf
LexA-p75NGFR(338-396)	+	•
LexA-p75NGFR(365-396)	+	1
LexA-Fas	++	1
LexA-Ras <sup>V12</sup>		+
LexA-Lamin	•	

**FIG. 13A** 

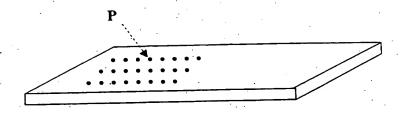
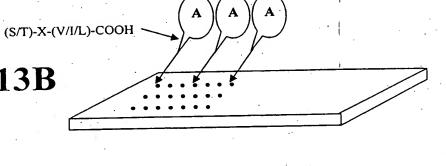
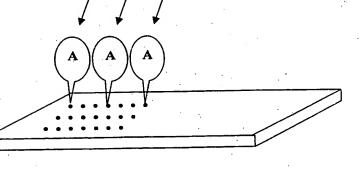


FIG. 13B







### FIG. 14A Plain-glass slide FIG. 14B 3D gel pad chip FIG. 14C Microwell chip